

Attorney Docket No.: ACY-33377

**Patent** 

### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Metcalf et al.

Serial No.: 09/674,368

(entry into U.S. National Stage from

PCT/US99/09486

Filed: April 29, 1999

(PCT filed January 29, 1998)

For: Vaccines Containing Recombinant Pilin

Against Neisseria Gonhorrhoeae or Neisseria

Meningitidis

Group Art Unit: TBD

Examiner: TBD

Atty. Dkt. No.: ACY-33377

### RESPONSE TO NOTIFICATION OF MISSING REQUIREMENTS UNDER 35 U.S.C. § 371

Commissioner for Patents
Box PCT
U.S. Patent and Trademark
Office
P.O. Box 2327
Arlington, Va 22202

**CERTIFICATION UNDER 37 CFR 1.10** 

I hereby certify that this paper and the documents referred to as enclosed therein are being deposited with the United States Postal Service on the date written below in an envelope as "Express Mail Post Office to Addressee" Mailing Label Number ET937267823US addressed to the Commissioner for Patents, Box PCT, U. S. Patent and Trademark Office, P.O. Box 2327, Arlington, VA 22202.

12/11/01

Date

Anand P. Iyer, Ph.D.

Sir:

In a "Notification of Missing Requirements under 35 U.S.C. §371", (the "Notification") mailed June 11, 2001, the Examiner stated that the Applicants had not submitted the required sequence listing pursuant to 37 C. F. R. §1.821 - §1.825. The Examiner also set a two-month time-period to comply with the Notification from the mailing date.

In response, Applicants submit herewith a substitute computer readable form of the Sequence Listing as required under 37 C.F.R.§1.821-§1.825, along with a copy of the

Notification. Applicants also submit that the sequence information provided in the

Specification is the same as the sequence listing of the enclosed substitute computer readable

form of the sequence listing. The Applicants, in accordance with 37 C.F.R. §1.821(g), also

attest that no new matter is included with this submission.

In addition, the Applicants have also submitted herewith a "Petition for Extension of

Time", for four months extension with the authorization to charge the extension fees to

Applicants' Deposit Account No. 01-1300. If any additional fee is required, the Patent Office

is authorized to charge the Applicants' Deposit Account No. 01-1300.

Respectfully submitted,

Oln M. Horch

Alan M. Gordon Reg. No. 30,637

Attorney for Applicants

American Home Products Corp. Five Giralda Farms Madison, NJ 07940 (845) 602-2189

Date:

12/11/01



UNITED STATES PATENT AND TRADEMARK OFFICE Patent Law Department Pearl River, NY

Commissioner for Patents, Box PCT United States Patent and Trademark Office Washington, D.C. 20231

U.S APPLICATION NO.	FIRST NAMED APPLICANT	<u> </u>		ATTY. DOCKET NO.
09/674368	METCALF	T	INTERNATIONAL AP	33377-00
ALAN M GORDON PATENT LAW DEPARTMENT AMERICAN HOME PRODUCT			PCT/US99	9/09486
ONE CAMPUS DRIVE PARSIPPANY, NJ 07054	and the second s		APR 99	JUN 2001
NOTIFICATION OF MIS	SSING REQUIREMENTS UND	. •	DATE MAILED:	

	Car	(O) DATE MAILCED.	•
NOTIFICATION OF MISSING REQUESTATES DESIGNATED		S UNDER 35 U.S.C. 371 IN THI ED OFFICE (DO/EO/US)	E UNITED
1. The following items have been submitted by the Office as	1.494)   an El   Indication of Translation     Translation   Other:   Other:   tion Report in En     total Preliminary	lected Office (37 CFR 1.495): of Small Entity Status. I of the international application into Engli I of Article 19 amendments into English. I glish and its Annexes, if any. Examination Report into English.	sh.
2. Applicant has requested early processing under the indicated items in paragraph 3 below. The Basis prior to 20 or 30 months from the priority date to available U.S. Basic National Fee.	c National Fee an void abandonmen	id the copy of the international application	d items and/or must be filed
Translation.  b. Processing fee for providing the transpropriate 20 or 30 months from c. Oath or declaration of the inventors, the application (preferably by the surcharge will be required if substance.  The current oath or declaration declaration declaration of the indicated on the attached PCT/DC d. Surcharge for providing the oath or priority date (37 CFR 1.492(e)).	nglish. A process O months from the e for the reasons aslation of the app a the priority date in compliance w International app mitted later than to coes not comply w O/EO/917. declaration later large entity additional claim	sing fee will be required if submitted e priority date. indicated on the attached Notice of Defect plication and/or the Annexes later than the (37 CFR 1.492(f)). With 37 CFR 1.497(a) and (b), properly idealization number and international filing dathe appropriate 20 or 30 months from the prith 37 CFR 1.497(a) and (b) for the reason than the appropriate 20 or 30 months from small entity, including any required multiples or cancel the additional claims for which the priority date of the same of	ntifying tte). A priority ns a the ple dependent nich fees are
ALL OF THE ITEMS SET FORTH IN 3(a)-3(d) MONTHS FROM THE DATE OF THIS NOTIC THE PRIORITY DATE FOR THE APPLICATI RESPOND WILL RESULT IN ABANDONMEN	CE OR BY 22 OF ON, WHICHEY	R 32 MONTHS (where 37 CFR 1.495 a)	plies) FROM
The time period set above may be extended by filin 1.136(a).		· · · · · · · · · · · · · · · · · · ·	
6: If box 3a or 3c is checked, a translation of the Annexes will be cancelled. A processing fee will b 7.  The Article 19 amendments are cancelled sin or 30 (37 CFR 1.495(d)) months from the priority	e required if subs see a translation w	mitted later than 20 or 30 months from the	priority date.
Applicant is reminded that any communication to the address given in the heading and include the U.S. a			ed to the
Enclosed: PCT/DO/EO/917 PTO-875	MUST be rosuse of Defective	Deborah D. Williams	JUN 1 3 2001
FORM PCT/DO/EO/905 (March: 2001)	•	71 iephone: 703-305-3744	ANNETTE CROMER

BEST AVAILABLE

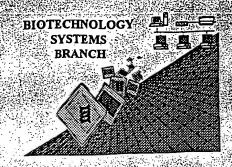
POTU B USE					Washington, D.C.
U.S. APPLICATION NO.		FIRS	T NAMED APPLICANT		ATTY, DOCKET NO.
09/674	368	ME	TCALF	T	33377-00
				INTERNATI	ONAL APPLICATION NO.
ALAN M GORDON PATENT LAW DE		20		PCT.	/US99/09486
		S CORPORATION		I.A. FILING DAT	PRIORITY DATE
ONE CAMPUS DR				29 APR 99	
PARSIFFAIT, INS	07054				_
				DATE MAIL	mil 1 Jun 2001
NOTIFICATION CONTAIN	ON TO CO	MPLY WITH REC LEOTIDE SEQUE	QUIREMENTS NCE AND/OF	S FOR PATENT R AMINO ACII	F APPLICATIONS D SEQUENCE
			LOSURES		
America. The ite	ms indicate	ers under 35 U.S.C. d below, however, a void abandonment is	re missing. Th	ne period within	in the United States of which to correct the Notification.
The nucleotide an with the requirem reason(s):	d/or amino ents for suc	acid sequence disclo h a disclosure as set	sure contained forth in 37 CF	in this application R 1.821-1.825 f	on does not comply or the following
· The ap	plication fa	ils to comply with th	ne requirements	of 37 CFR 1.83	21_1 825
		oes not contain, a "S			
		er copy or compact of			
A copy	of the "Se	quence Listing" in c	omputer readah	ole format has no	t been submitted as
require	d by 37 CF	R 1.821(e).			
A copy	of the "Se	quence Listing" in c	omputer readab	ole form has been	1 submitted. The
37 CFF	of the com R 1.822 and ce Listing."	or 1.832, as indicat	, however, doe ed on the attacl	s not comply with hed marked-up c	th the requirements of opy of the "Raw
			een filed with	this application l	nas been found to be
damag substit	ed and/or u ute compute	nreadable as indicate or readable form must compact disc of the	ed on the attach	ed CRF Diskette as required by 3	e Problem Report. A
		e form of the "Seque			
Other:		<u> </u>			
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APPLICANT MU	IST PROVI	DE:			
		itute computer reada	ble form (CRF	) of the "Sequen	ce Listing "
					Listing," as well as an
		ing its entry into the			sioning, as well as all
A state	ment that th	ne contents of the pa	per or compact	disc and the cor	mputer readable form
are the	same and,	where applicable, in 1.821(g), 1.825(b)	clude no new n		
FOR QUESTION CALL:	S REGARE	ING COMPLIANC	E WITH THES	SE REQUIREM	ENTS, PLEASE
(703) 308	3-4216, for	Rules interpretation,			
(703) 308	3-4212, for	CRF submission hel PatentIn software he	p,		
	•				

Deborah D. Williams

FORM PCT/DO/EO/920 (March 2001)

BEST AVAILABLE COPY

# RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

PUT/09

Source:

c: 3/01/2001

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: <a href="mailto:patin21help@uspto.gov">patin21help@uspto.gov</a> or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: <a href="mailto:patin3help@uspto.gov">patin3help@uspto.gov</a> or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

#### Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

# Raw Sequence Listing Error Summary

i	ERROR DETECTED	SUGGESTED	CORRECTION	٠	SERIAL NUMBER:	9/674,368
ATTA	I- NEW RUI ES CASES: F	PLEASE DISREGA	RD ENGLISH "ALPH	A" HEADERS, W	HICH WERE INSERTED E	BY PTO SOFTWARE
, ^Y"	Wrapped Nucleics	The number/text	at the end of each line	"wrapped" down l	o the next line.	
	_ ***apped ***do.e.do	This may occur if	f your file was retrieved	d in a word proces	sor after creating it.	
		Please adjust yo	our right margin to .3, a	s this will prevent	"wrapping".	
2	Wrapped Aminos	The amino acid n	number/text at the end	of each line Wrap	ped a down to the next line.	•
·	. Wrapped Aminos	This may occur i	if your file was retrieve	d in a word proces	sor after creating it.	
	*	Place adjust you	ur right margin to .3, a	s this will prevent	"wrapping".	
3	Incorrect Line Length	The rules require	that a line not exceed	72 characters in le	ength. This includes space	es. •
4	Misaligned Amino Acid	The numbering u	nder each 5lh amino a	icid is mişaligned.	This may be caused by the	use of tabs
1	Numbering	between the number	bering. It is recommen	nded to delete any l	labs and use spacing between	een the numbers.
/ ا ء	Non-ASCII	This file was not s	saved in ASCIL (DOS)	text, as required b	y the Sequence Rules.	•
3 <u>U</u>	Non-ASCII	Please ensure vor	ur subsequent submis	sion is saved in A	SCII text so that it can be p	rocessed.
					-	· · ·
6	Variable Length	Sequence(s)	_ contain n's or Xaa's	which represented	more than one residue.	
	-	As per the rules, e	each n or Xaa can only	represent a singl	e residue.	
		Please present the	e maximum number of	reach residue nav	ing variable length and	
		indicate in the (ix)	) feature section that s	some may be missi	ng.	•
7	Patentin ver. 2.0 "bug"	A "bug" in Palentl	In version 2.0 has caus	sed the <220>-<22	23> section to be missing fo	rom amino acid
		sequence(s)	. Normally,	Patentin would at	itomatically generate this se	ection from the
		previously coded	nucleic acid sequence	e. Please manually	y copy the relevant <220>	<223> section
		to the subsequen	it amino acid sequence	e. This applies p	rimarily to the mandatory	y <220>-<223>
		sections for Artif	ficial or Unknown se	quences.		
,	Skipped Sequences	Sequence(s)	missing If intentiona	l please use the fe	ollowing formal for each ski	ipped sequence:
' ——	(OLD RULES)		N FOR SEQ ID NO:X		•	
	(OLD NOLLO)	(i) SEQUENCE C	HARACTERISTICS:	(Do not insert any l	headings under "SEQUEN	CE CHARACTERISTICS")
			DESCRIPTION:SEQ			
			s intentionally skippe			
		Please also adjust	t the "(iii) NUMBER O	F SEQUENCES:"	response to include the sk	ipped sequence(s).
					-	
·	Skipped Sequences	Sequence(s)		i, piease use the it	ollowing format for each ski	pped sequence.
	(NEW RULES)	<210> sequence				
		₹400> sequence	la number		•	
		000			•	
)	Use of n's or Xaa's	Use of n's and/or ?	Xaa's have been delec	ted in the Sequen	ce Listing.	
	(NEW RULES)	Use of <220> to <	223> is MANDATOR'	Y if n's or Xaa's are	e present.	
	(	In <220> to <223>	section, please expla	in location of n or	Xaa, and which residue n	or Xaa represents.
	Use of a313's Occasion	Somunaco(s)	are missing this r	mandatory field of i	ts response.	
	Use of <213>Organism (NEW RULES)	Sequence(s)	are massing mas i	indication y management	-	
		•	•	•	<u> </u>	·-
	Use of <220>Feature	Sequence(s)	are missing the <220	>Feature and asso	ociated headings.	
	(NEW RULES)				ISM is "Artificial" or "Unkn	own"
		Please explain so	ource of genetic mat	erial in <220> lo	<223> section.	( 0
/	^	(See "Federal	Register," 6/01/9	98, Vol. 63, No	. 104, pp. 29631-32)	(Sec. 1.823 of new Ru
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<u></u>	Palentin ver. 2.0 oug	Tital Take up not use	ssing mandaloty num	eric identifiers and	responses (as indicated or	raw sequence listing).
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		picase usi		•	• • •	

Does Not Comply

PC'

Corrected Diskette Needer Input Set : A:\PTO.txt Output Set: N:\CRF3\03072001\1674368.raw 5 <110> APPLICANT: American Cyanamid Company 7 <120> TITLE OF INVENTION: Vaccines Containing Recombinant Pilin Against Neisseria Gonorrhoeae or Neisseria Meningitidis sel dem 5 on End 11 <130> FILE REFERENCE: 33377-00/PCT 13 <140> CURRENT APPLICATION NUMBER: US/09/674,368 14 <141> CURRENT FILING DATE: 2000-10-27 Summary Sheet 16 <160> NUMBER OF SEQ ID NOS: 24 18 <170> SOFTWARE: PatentIn Ver. 2.0 ERRORED SEQUENCES 299 <210> SEQ ID NO: 13 move up - format enow (see item I on Even Summany Sheet) 300 <211> LENGTH: 31 301 <212> TYPE: DNA 303 <213> ORGANISM: Neisseria Meningitidis Class I 305 <400> SEQUENCE: 13 E--> 306 ccggcgcgtc tctcacggcg aatggcccgg c 307 396 <210> SEQ ID NO: 24 397 <211> LENGTH: 22 398 <212> TYPE: PRT 399 <213> ORGANISM: Neisseria gonorrhoeae 404 <400> SEQUENCE: 24 405 Glu Ala Ile Leu Leu Ala Glu Gly Gln Lys Ser Ala Val Thr Glu Tyr 10 406 1 408 Tyr Leu Asn His Gly Lys 20 410 E--> 441/1 E--> 443/8 delete at end I file ACY-33377-00 see follning pages for more error

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/674,368

DATE: 03/07/2001

TIME: 11:37:48

VERIFICATION SUMMARY DATE: 03/07/2001 PATENT APPLICATION: US/09/674,368 TIME: 11:37:50

Input Set : A:\PTO.txt

Output Set: N:\CRF3\03072001\1674368.raw

L:13 M:270 C: Current Application Number differs, Replaced Application Number L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:60 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1 L:64 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1 L:68 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1 L:72 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1 L:76 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1 L:80 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1 L:81 M:258 W: Mandatory Feature missing, <220> FEATURE: L:87 M:258 W: Mandatory Feature missing, <220> FEATURE: L:194 M:258 W: Mandatory Feature missing, <220> FEATURE: L:194 M:258 W: Mandatory Feature missing, <220> FEATURE: L:194 M:258 W: Mandatory Feature missing, <220> CTHER INFORMATION: L:306 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:31 SEQ:13

L:441 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:24 M:332 Repeated in SeqNo=24

4

RAW SEQUENCE LISTING DATE: 03/14/2001 PATENT APPLICATION: US/09/674,368 TIME: 12:10:34

Input Set : A:\Pto.amc

Output Set: N:\CRF3\03142001\1674368.raw

	142	1				5					10					15		0.0				
	144	atc	gcc	atc	gtc	ggt	atc	ttg	gca	gcc	gtc	gcc	ctg	ccc	gca	tac	caa	96				
	145	Ile	Ala	Ile			Ile	Leu	Ala			Ala	Leu	Pro	Ala	Tyr	GIn					
	146				20					25					30							
	148	gac	tac	acc	gcg	cgc	gcc	caa	atg	tcc	gaa	gcc	ctg	act	ttg	gca	gaa	144				
	149	Asp	Tyr	Thr	Ala	Arg	Ala	Gln		Ser	Glu	Ala	Leu	Thr	Leu	Ala	Glu		•			
	150			35					40					45				100				
	152	ggt	caa	aaa	tcc	gca	gtg	atc	gag	tat	tat	tcc	gac	aac	ggc	aca	ttc	192				
	153	Gly	Gln	Lys	Ser	Ala	Val		Glu	Tyr	Tyr	Ser		Asn	Gly	Thr	Pne					
	154		50					55					60					240				
	156	ccg	aac	agc	aat	act	tcc	gca	ggt	att	gct	gcc	tct	aac	gag	att	aaa -	240				
	157	Pro	Asn	Ser	Asn	Thr		Ala	Gly	Ile	Ala		Ser	Asn	Glu	He						
	158	65					70					75					80	200				
	160	ggt	aag	tat	gtg	gca	tcg	gtt	aag	gtt	gaa	ggt	aat	gcc	tct	gtt	gct	288				
	161	Gly	Lys	Tyr	Val		Ser	Val	Lys	Val			Asn	Ala	Ser		Ala					
	162					85					90					95						
	164	tct	att	acc	gct	acc	atg	aac	tct	agt	aat	gtg	aat	aag	gac	atc	aaa	336				
	165	Ser	Ile	Thr	Ala	Thr	Met	Asn	Ser		Asn	Val	Asn	Lys	Asp	He	Lys					
	166				100					105					110							
	169	ggt	aaa	acc	ttg	gta	ctc	gtc	ggc	aaa	caa	aac	tcc	ggt	tcg	gta	aaa	384				
	170	Gly	Lys	Thr	Leu	Val	Leu	Val	Gly	Lys	Gln	Asn	Ser	Gly	Ser	Val	Lys					
•	171			115					120					125								
	173	tgg	ttc	tgc	gga	cag	ccg	gtt	acg	cgc	gac	aac	gcc	gac	aac	gac	gac	432				
	174	Trp	Phe	Cys	Gly	Gln	Pro	Val	Thr	Arg	Asp	Asn		Asp	Asn	Asp	Asp					
	175		130					135					140									
	177	gtc	aaa	gac	gcc	ggc	aac	aac	ggc	atc	gaa	acc	aag	cac	ctg	ccg	tca	480				
	178	Val	Lys	Asp	Ala	Gly	Asn	Asn	Gly	Ile	Glu		Lys	His	Leu	Pro	Ser					
	179						150					155					160					
	181	acc	tgc	cgc	gat	acg	tca	tct	gat	gcc	aaa							510				
	182	Thr	Cys	Arg	Asp	Thr	Ser	Ser	Asp	Ala	Lys						•					
	183					165					170											
	186	<210	)> SE	EQ [[	ON C	: 4																
	187	<211	> LE	ENGT	1: 17	70														•		_
		<212																		٠. د	1	17
	189	521g	> OF	RGANI	SM:	Arti	fici	al S	Seque	ence				$\sim$	1	1 2	1 0	121 -	sle	Nes	2 /	3 Heet
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W>	19	<223	ro k	HER	INFO	ORMAT	NOI?	:		_			/						$\mathcal{O}_{n}$	5/100	<b>^</b> .	
	191	<u> </u>	> SE	EQUE	ICE:	4													071	000	•	
	192	Met	Glu	Ala	Ile	Gln	Lys	Gly	Phe	Thr	Leu	Ile	Glu	Leu	Met	Ile	Val		1			11 +
	193					5					10					15			Aus	mar	u, /	Her
	195	Ile	Ala	Ile	Val	Gly	Ile	Leu	Ala	Ala	Val	Ala	Leu	Pro	Ala	Tyr	Gln		100		1	
	196				20					25					30							
	198	Asp	Tyr	Thr	Ala	Arg	Ala	Gln	Met	Ser	Glu	Ala	Leu	Thr	Leu	Ala	Glu					
	199			35					40					45								
	201	Gly	Gln	Lys	Ser	Ala	Val	Ile	Glu	Tyr	Tyr	Ser	Asp	Asn	Gly	Thr	Phe					
	202	_	50					55					60									
	204	Pro	Asn	Ser	Asn	Thr	Ser	Ala	Gly	Ile	Ala	Ala	Ser	Asn	Glu	Ile	Lys					
	205						70					75					80					
	207	Glv	Lys	Tyr	Val	Ala	Ser	Val	Lys	Val	Glu	Gly	Asn	Ala	Ser	Val	Ala					
		•	-	-																		

DATE: 03/14/2001

PATENT APPLICATION: US/09/674,368 TIME: 12:10:34 Input Set : A:\Pto.amc Output Set: N:\CRF3\03142001\1674368.raw 71 gcc aag acc gcc gac aac atc aac acc aag cac ctg ccg tca acc tgc 72 Ala Lys Thr Ala Asp Asn Ile Asn Thr Lys His Leu Pro Ser Thr Cys 150 155 75 cgc gac gca agt gat gcc agc taa 504 76 Arg Asp Ala Ser Asp Ala Ser > 77 79 <210> SEQ ID NO: 2 80 <211> LENGTH: 167 whet I le Val

So Ala Tyr Gln

30

Su Leu Ala Glu

5

5 Gly Glu Trp

Men Ala Tyr

Men Ala Glu

Men Ala Glu 81 <212> TYPE: PRT 82 ORGANISM: Artificial Sequence W--> 84 <220> FEATURE: 84 (223) OTHER INFORMATION: 84 (400) SEQUENCE: 2 W--> 84 85 Met Asp Thr Leu Gln Lys Gly Phe Thr Leu Ile Glu Leu Met Ile Val 86 1 10 88 Ile Ala Ile Val Gly Ile Leu Ala Ala Val Ala Leu Pro Ala Tyr Gln 25 92 Asp Tyr Thr Ala Arg Ala Gln Val Ser Glu Ala Ile Leu Leu Ala Glu 35 40 96 Gly Gln Lys Ser Ala Val Thr Glu Tyr Tyr Leu Asn His Gly Glu Trp 50 55 60 100 Pro Gly Asn Asn Thr Ser Ala Gly Val Ala Ser Ser Ser Thr Ile Lys 101 65 104 Gly Lys Tyr Val Lys Glu Val Thr Val Ala Asn Gly Val Ile Thr Ala 108 Thr Met Leu Ser Ser Gly Val Asn Lys Glu Ile Gln Gly Lys Lys Leu 100 105 112 Ser Leu Trp Ala Lys Arg Gln Asp Gly Ser Val Lys Trp Phe Cys Gly 115 120 115 Gln Pro Val Thr Arg Thr Asp Ala Lys Ala Asp Thr Val Ala Ala Ala 135 140 118 Ala Lys Thr Ala Asp Asn Ile Asn Thr Lys His Leu Pro Ser Thr Cys 119 145 150 155 121 Arg Asp Ala Ser Asp Ala Ser 122 165 125 <210> SEQ ID NO: 3 126 <211> LENGTH: 510 127 <212> TYPE: DNA 128 <213> ORGANISM: Artificial Sequence 130 <220> FEATURE: 131 <223> OTHER INFORMATION: Description of Artificial Sequence: Chimeric of Neisseria Meningitidis Class II and Neisseria 133 Gonorrhoeae 135 <220> FEATURE: 136 <221> NAME/KEY: CDS 137 <222> LOCATION: (1)..(510) 139 <400> SEQUENCE: 3 140 aty gaa gca atc caa aaa ggt ttc acc ctg atc gag ctg atg atc ytc 141 Met Glu Ala Ile Gln Lys Gly Phe Thr Leu Ile Glu Leu Met Ile Val

RAW SEQUENCE LISTING

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